

VALIDATION RESULTS FOR METHOD NO. MP-NIPM-MA						
		MEAN	RELATIVE STD. DEV.(%)	RANGE	SEE (conc)	
ANALYTE	NNAL					
MATRIX	Human urine					
DATE	14.Sept.01					
LLOQ	0.6 ng/100 mL					
	Accuracy 71.7%					
	RSD 35.0%					
STANDARD ERROR OF ESTIMATE OF QC SAMPLES (concentration units) (SEE = S.D./SQRT (# of determinations))						
	0.6 ng/100 mL	Low QC			0.18	
	75 ng/100 mL	Med QC			3.62	
	200 ng/100 mL	High QC			9.58	
DESCRIPTIVE INFORMATION ON INSTRUMENT RESPONSE						
Function (Linear, Quadratic, etc.)	Linear					
Slope (mean)	18.53					
Intercept (mean)	0.08					
Other (mean regression value)	0.99679(1/y weighted)					
RECOVERY (as a %)						
High QC in (matrix)=	200 ng/100 mL	71.4%	Calculated as a % of the concentration observed for derivatized urine standards			
Med QC in (matrix)=	75 ng/100 mL	81.8%				
Low QC in (matrix)=	1.8 ng/100 mL	116.1%				
Results with different sample amounts (results of QC evaluations) intra-assay						
LLOQ QC=	0.6 ng/100 mL	0.43	35.0%	0.24-0.58	0.06	
Low QC=	1.8 ng/100 mL	1.34	6.6%	1.23-1.47	0.04	
Med QC=	75 ng/100 mL	70.03	7.5%	64.85-77.56	2.14	
High QC=	200 ng/100 mL	179.46	8.4%	158.48-196.94	6.12	
Results with different sample amounts (results of QC evaluations) inter-assay						
LLOQ QC=	n/a	n/a	n/a	n/a	n/a	
Low QC=	1.8 ng/100 mL	1.8	35.0%	1.23-3.83	0.16	
Med QC=	75 ng/100 mL	84.2	18.2%	64.85-120.54	3.62	
High QC=	200 ng/100 mL	215.6	18.9%	158.48-309.65	9.58	
STABILITY (assuming any degradation is linear):						
a) Room Temperature in Matrix (+/- %/24 hrs)	-18.4	% maximum				
b) Freeze/thaw (+/- %/3 cycles)	-19.7	% maximum				
c) Injector (+/- %/24 hrs)	-31.5	% maximum				